

Untitled
GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: June 19, 2005, 21:40:17 ; Search time 24770 Seconds
(without alignments)
12052.439 Million cell updates/sec

Title: US-09-242-772-116

Perfect score: 7313

Sequence: 1 ggcagcgcatacactacaat.....tatgaataaaatctcgtgcc 7313

Scoring table: OLIGO_NUC

Gapop 60.0 , Gapext 60.0

Searched: 45554873 seqs, 20411521753 residues

Word size : 0

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : Pending_Patents_NA_Main:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	% Length	DB	ID	Description
1	7313	100.0	7313	18	US-09-242-772-116	Sequence 116, App
2	7313	100.0	7313	41	US-09-948-941-30	Sequence 30, Appl
3	7313	100.0	7313	41	US-09-949-003C-307	Sequence 307, App
4	7313	100.0	7313	64	US-10-896-891-30	Sequence 30, Appl
5	7313	100.0	7313	129	US-60-592-408-133	Sequence 133, App
6	3910	53.5	7107	41	US-09-948-941-221	Sequence 221, App
7	3910	53.5	7107	41	US-09-949-003C-1603	Sequence 1603, Ap
8	3910	53.5	7107	49	US-10-170-235-35981	Sequence 35981, A
9	3910	53.5	7107	64	US-10-896-891-221	Sequence 221, App
10	3767	51.5	7312	86	US-60-172-373-5545	Sequence 5545, Ap
11	3350	45.8	22554	41	US-09-948-941-554	Sequence 554, App
12	3350	45.8	22554	41	US-09-948-941-745	Sequence 745, App
13	3350	45.8	22554	41	US-09-949-003C-3997	Sequence 3997, Ap
14	3350	45.8	22554	41	US-09-949-003C-5293	Sequence 5293, Ap
15	3350	45.8	22554	64	US-10-896-891-554	Sequence 554, App
16	3350	45.8	22554	64	US-10-896-891-745	Sequence 745, App
17	1037	14.2	1210	34	US-09-760-498-104	Sequence 104, App
18	1037	14.2	1210	48	US-10-143-788-104	Sequence 104, App
c 19	601	8.2	601	41	US-09-948-941-1506	Sequence 1506, Ap
c 20	601	8.2	601	41	US-09-948-941-10216	Sequence 10216, A
c 21	601	8.2	601	41	US-09-949-003C-14347	Sequence 14347, A

Untitled

c 22	601	8.2	601	41	US-09-949-003C-62838	Sequence 62838, A
c 23	601	8.2	601	64	US-10-896-891-1506	Sequence 1506, Ap
c 24	601	8.2	601	64	US-10-896-891-10216	Sequence 10216, A
c 25	550	7.5	601	41	US-09-947-907-9408	Sequence 9408, Ap
c 26	550	7.5	601	41	US-09-948-941-1503	Sequence 1503, Ap
c 27	550	7.5	601	41	US-09-948-941-10213	Sequence 10213, A
c 28	550	7.5	601	41	US-09-949-003C-14344	Sequence 14344, A
c 29	550	7.5	601	41	US-09-949-003C-62835	Sequence 62835, A
c 30	550	7.5	601	64	US-10-896-891-1503	Sequence 1503, Ap
c 31	550	7.5	601	64	US-10-896-891-10213	Sequence 10213, A
c 32	432	5.9	601	41	US-09-947-907-9407	Sequence 9407, Ap
c 33	432	5.9	601	41	US-09-948-941-1505	Sequence 1505, Ap
c 34	432	5.9	601	41	US-09-948-941-10215	Sequence 10215, A
c 35	432	5.9	601	41	US-09-949-003C-14346	Sequence 14346, A
c 36	432	5.9	601	41	US-09-949-003C-62837	Sequence 62837, A
c 37	432	5.9	601	64	US-10-896-891-1505	Sequence 1505, Ap
c 38	432	5.9	601	64	US-10-896-891-10215	Sequence 10215, A
39	430	5.9	566	32	US-09-716-990-1050	Sequence 1050, Ap
c 40	423	5.8	599	29	US-09-644-869-8143	Sequence 8143, Ap
41	414	5.7	415	21	US-09-362-510-13920	Sequence 13920, A
42	414	5.7	415	21	US-09-362-510A-13920	Sequence 13920, A
43	414	5.7	415	39	US-09-904-013-13920	Sequence 13920, A
c 44	374	5.1	522	24	US-09-535-897-2429	Sequence 2429, Ap
c 45	360	4.9	601	41	US-09-948-941-1504	Sequence 1504, Ap

Untitled

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: June 19, 2005, 21:12:51 ; Search time 1074 Seconds
(without alignments)
11141.612 Million cell updates/sec

Title: US-09-242-772-116

Perfect score: 7313

Sequence: 1 ggtagcgcatacactacaat.....tatgaataaaatctcggtgcc 7313

Scoring table: OLIGO_NUC

Gapop 60.0 , Gapext 60.0

Searched: 1202784 seqs, 818138359 residues

word size : 0

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : Issued_Patents_NA:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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3	35	0.5	660	4	US-09-270-767-22908		Sequence 22908, A
c 4	34	0.5	601	4	US-09-949-016-28962		Sequence 28962, A
c 5	34	0.5	601	4	US-09-949-016-125909		Sequence 125909,
c 6	34	0.5	115963	4	US-09-949-016-12298		Sequence 12298, A
c 7	34	0.5	144322	4	US-09-949-016-15316		Sequence 15316, A
c 8	33	0.5	601	4	US-09-949-016-115142		Sequence 115142,
c 9	33	0.5	2398	4	US-09-687-637B-21		Sequence 21, Appl
10	33	0.5	11730	4	US-09-949-016-14991		Sequence 14991, A
c 11	33	0.5	49416	4	US-09-949-016-15234		Sequence 15234, A
c 12	33	0.5	93364	4	US-09-949-016-14890		Sequence 14890, A
13	33	0.5	283538	4	US-09-949-016-13506		Sequence 13506, A
14	32	0.4	1118	4	US-09-439-554-19		Sequence 19, Appl
15	32	0.4	2075	4	US-09-461-325-130		Sequence 130, App
16	32	0.4	2075	4	US-10-012-542-130		Sequence 130, App
17	32	0.4	2075	4	US-10-115-123-130		Sequence 130, App

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c 18	32	0.4	100550	4	US-09-949-016-11835	Sequence 11835,	A
c 19	32	0.4	100551	4	US-09-949-016-16207	Sequence 16207,	A
20	32	0.4	462589	4	US-09-949-016-12900	Sequence 12900,	A
21	32	0.4	476044	4	US-09-949-016-12412	Sequence 12412,	A
c 22	32	0.4	678533	4	US-09-949-016-14577	Sequence 14577,	A
c 23	32	0.4	678533	4	US-09-949-016-14578	Sequence 14578,	A
24	31	0.4	286	4	US-10-021-338A-35	Sequence 35,	Appl
c 25	31	0.4	321	2	US-08-520-678A-23	Sequence 23,	Appl
c 26	31	0.4	321	3	US-08-897-126-23	Sequence 23,	Appl
c 27	31	0.4	601	4	US-09-949-016-38299	Sequence 38299,	A
28	31	0.4	601	4	US-09-949-016-53527	Sequence 53527,	A
29	31	0.4	601	4	US-09-949-016-53537	Sequence 53537,	A
30	31	0.4	601	4	US-09-949-016-60222	Sequence 60222,	A
c 31	31	0.4	601	4	US-09-949-016-71451	Sequence 71451,	A
32	31	0.4	601	4	US-09-949-016-82033	Sequence 82033,	A
33	31	0.4	601	4	US-09-949-016-82038	Sequence 82038,	A
c 34	31	0.4	601	4	US-09-949-016-137776	Sequence 137776,	
c 35	31	0.4	601	4	US-09-949-016-146550	Sequence 146550,	
c 36	31	0.4	601	4	US-09-949-016-146551	Sequence 146551,	
c 37	31	0.4	601	4	US-09-949-016-169981	Sequence 169981,	
c 38	31	0.4	601	4	US-09-949-016-189744	Sequence 189744,	
39	31	0.4	997	4	US-09-523-686-1	Sequence 1,	Appl
40	31	0.4	1847	4	US-09-904-615-41	Sequence 41,	Appl
41	31	0.4	2594	4	US-09-602-472A-7	Sequence 7,	Appl
42	31	0.4	2793	4	US-09-602-472A-9	Sequence 9,	Appl
43	31	0.4	3829	2	US-08-631-097-8	Sequence 8,	Appl
44	31	0.4	3829	3	US-08-810-712-6	Sequence 6,	Appl
c 45	31	0.4	4741	1	US-07-695-472B-4	Sequence 4,	Appl